Figure 1

M R S																																				AG 90
L E D L L E G S Q Q L E D F A Y A Y P E R N R V F G G K A H GACGACA-GGGTTAACTATCTCTCACGAGGACTGAAGAACA-CTGGCTACTATGATGTCTACAAGCAGCCTCAGGTGCACCTGTGGAGCAAT 27 D D T V N Y L Y E E L K K T G Y Y D V Y K O P Q V H L W S N GCCGACCAGAGCGCTCAAGGTGGGCGATGAGGAAGACCATGACCTACGAGTCCCAGGCTCCAGGTGCACCTGGAGCAACT GCCGACCAGACGCTCAAGGTGGGGGAATACCAGGAGCGAAGACCATGACCTACCAGCCTCGAGGTCACCGCCGATGTACC A D Q T L K V G D E E I E A K T M T Y \$ P S V E V T A D V A GTCGTCAAGAACCTGGGATGCAGCGAGGGGGGATTACCCATCCCATGTCGAGGGCAAAGGTCGCCCTGATCAAGCGTGGAGAATGCCCGTTC 45 V V K N L G C S E A O Y P S D V E G K V A L I K R G E C P F GGCGACAAGTCGGTTCTCGCTGCCAAAGCCAAGGCCGAGGCGGCGTTCCATGTCGAGGGGGAACGTTCGCCCTGATCAAGCATGCTGGCGGGCAACGTTCGGCCGAGACCTTGGC 544 G D K S V L A A K A A A A S I V Y N N V A G S H A G T L G GCGGGCGCAGAGTGATAAGGGAACCGTATTCGGCCATTGTCGGTATCAACTTGTGGCAGAACGATCAACCTTGGCTAGGCCGAGACCGAACCGTACAAGCTTGGTCAGGCTGAGCTAGAACAGCTTGCAGACTGAGCTAGAGCTAGAACCTGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAACGTATCAAGCTTGGTCAGAGCTAGAACAGCTTGGTCAGAGCTAGAACAGCTTGGTCAGAGCTAGAACAGCTAGAACAGCTAGAACAGCTAGAACAGCTAGAACAGCTAGAACAGAGAAACCGTAGAACAGAGAAACCGTATAACGTTGTGGGAACAGAGAGAG	<u>M</u>	R		<u> </u>		L	W	<u> </u>	S				s	G			_	<u>A</u>	G	R		Α	L	V		5 1	_	0	Ε	F	٩	<u> </u>	<u> </u>	0	1_	<u>a</u>
GACGACAGGTTAACTATCTCTACGAGGAGCTGAAGAAGACTGGCTACTATGATGTCTACAAGCAGCCTCAGGTGCACCTGGAGCAAT 27 O D T V N Y L Y E E L X X T G Y Y O V Y K Q P Q V H L W S N GCGACCAGACGCTCAAGGTGGAGGATGAGCCTAGGTGAAGCAGCTCCAAGGTGAGCCTAAGGTGAGCCTAAGGTGAGCCTAAGGTGAGCCTAAGGTGAGCCTAAGGTGAAGCATTAAGCAGTGAAGAACCTGAGAGCCTAAGGTGAAGAACCTGAGAGTGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAAGCCAAGGCCGAGAAGCTCAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTGGAAGAACCTAGGAAGCCAAGGCCAAGGCCGAGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCAAGGCCACCTTGGC S4C V N N N N N N N N N N N N N N N N N N	TT	GGA	AGA	TC	TGC	TGG	AA	GGA	TC	CCA	ACA	AGC	TT	GAG	GA	CT1	CG	CCI	ΓΑΤ	GC	CT	ACC	cc	GA	GCG	CAA	TC	GCG	TC	TT.	rgg	TG	STA	AAG	ccc	AC 180
D	L	Ξ	0		L_	L	Ε	G	S	(2 (<u></u>	L_	Ε	0	F	:	A	Y	Α	_ `	Y	۾	ε	R	! !		R	٧	F	G	(3	K	Α	Н
D	GA	CGA	CAC	GG.	TTA.	ACT	AT	СТС	TA	CGA	GGA	GC	TG	AAG	AA	GAC	:TG	GCT	TAC	TA	TGA	ATG	TC	TAČ	AA	GCA	GC	ITC	AG	GTO	CA	сст	GT	GGA	GC A	AT 270
A D Q T L K V G D E E I E A K T M T Y S P S V E V T A D V A GTCGTCAAGAACCTGGGATGCAGCGAGGCGAGGCGATTACCCATCCGATGTCGAGGGCAAGGTCGCCCTGATCAAGCGTGGAGAATGCCCGTTC 45: V V K N L G C S E A D Y P S D V E G K V A L I K R G E C P F GGCGACAAGTCGGTTCTCGCTGCCAAAGCCAAGGCCAGGCCTTCGATTGTCTTATAACAATGTGGCCGGATCCATGGCGGGCACCCTTGGC 540; G D K S V L A A K A K A A A S I V Y N N V A G S M A G T L G GCGGGCGCAGAGTGATAAGGGACCGTATTCGGGCATTGTCGGTTACCAGCTTGAGGAGAGGCTGAACCTTGCTGGAGAGTGATCAAGCTTGCTGAAGGCTGGA 630; A A Q S D K G P Y S A I V G I S L E D G D K L I K L A E A G CTCGGTATCTGTGGATCTGTGGGTGGATAGTAAGCAGGAGAACCGTACGACGTATAACGTTGTCGGCCAGAAGGCGGAACGGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACAGGCGGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACCGAACACGAACGCGAACCGAACACGAACACGAACACGAACACGAACACGAACACGAACACGAACACGAACACGAACCGAACACACACACACACACACACACACACACACACACACACA	0	Đ	r	1	v i	N	Y	L	Y	ε		. 1	L	K	K	Ţ	•	G	Y	Y	0	כ	٧	Y	K	C	1	9	α	٧	Н	L		W	S	N Z/G
A D Q T L K V G D E E I E A K T M T Y S P S V E V T A D V A GTCGTCAGAGACCTGGGATGCAGCGAGGCGAGGCGATTACCCATCCGATGTCGAGGGCAAGGTCGCCCTGATCAAGCGTGGAGAATGCCCGTTC 45: V V K N L G C S E A D Y P S D V E G K V A L I K R G E C P F GGCGACAAGTCGGTTCTCGCTGCCAAAGCCAAGGCCAGGCCTTCGATTGTCTTATAACAATGTGGCCGGATCCATGGCGGGCACCCTTGGC 540; G D K S V L A A K A K A A A S I V Y N N V A G S M A G T L G GCGGCGCAGAGTGATAAGGGACCGTATTCGGCCATTGTCGGTTATCAGCTTGGAGGATGGCCAGAAGCTGATCAAGCTTGCTGAGGCTGGA 630; A A O S D K G P Y S A I V G I S L E D G D K L I K L A E A G TCGGTATCTGTGGATCTGTGGGTGGATAGTAAGCAGGAGAACCGTACGACGTATAACGTTGTCGGCCAGAAGGCGGAACGGAACCGAACCGAACCGAACCAGACGAGAGGAG	GC	GA	CA:	GAC	GC.	TCA.	AGO	STG	GG	CGA	TGA	GGA	444	ATC	GA	GGC	GA.	AGA	CC.	ΑΤ	GAC	ст	'AC	AGT	·cc	CAG	CGI	.ce	Agr	370	'ΔΓ	rer	רכ	A TC	TAGG	r 360
V V K N L G C S E A O Y P S O V E G K V A L I K R G E C P F GGCGACAAGTCGGTTCTCGCTGCCAAAGCCCAAGGCCGGGCTTCGATTGTCTATAACAATGTGGCCGGATCCATGGCGGGCACCCTTGGC 546 G O K S V L A A K A K A A A S I V Y N N V A G S H A G T L G GCGGCGCAGAGTGATAAGGGACCGTATTCGGGCCATTGTCGGTATCAGCTTGGAGGATGGCCAGAAGCTGATCAAGCTTGCTGAGGCTGGA A A O S O K G P Y S A I V G I S L E O G O K L I K L A E A G TCGGTATCTGTGGATCTGTGGGTGGATAGAGCAGGAGGAGACGAAGCCGTACGACGTATAACGTTGTCGGCAGAAGCGAAGGGCGGAACCGAACGACGAAGGCGAACCGAAGCGAAGCGAAGGGCGGGAACCGAACCGAACCAACGACG	A	0	Q	1	Γι	٠.	K	٧	G	0	ε		Ξ	I	ε	A	. 1	K	Т	М	1	7	Y	Ś	P	s	1	,	E	v	T	A		ם סיים	٧ ,	300 4
V V K N L G C S E A O Y P S O V E G K V A L I K R G E C P F GGCGACAAGTCGGTTCTCGCTGCCAAAGCCCAAGGCCGGGCTTCGATTGTCTATAACAATGTGGCCGGATCCATGGCGGGCACCCTTGGC 546 G O K S V L A A K A K A A A S I V Y N N V A G S H A G T L G GCGGCGCAGAGTGATAAGGGACCGTATTCGGGCCATTGTCGGTATCAGCTTGGAGGATGGCCAGAAGCTGATCAAGCTTGCTGAGGCTGGA A A O S O K G P Y S A I V G I S L E O G O K L I K L A E A G TCGGTATCTGTGGATCTGTGGGTGGATAGAGCAGGAGGAGACGAAGCCGTACGACGTATAACGTTGTCGGCAGAAGCGAAGGGCGGAACCGAACGACGAAGGCGAACCGAAGCGAAGCGAAGGGCGGGAACCGAACCGAACCAACGACG	GTO	GTO	CAAC	GA <i>A</i>	CCI	rgge	GAT	rgc.	ΔGC	GΔ	GGC	GG A	. T T	- Δ C	cc.	۸۲۲	C 6 /	ATC	TC	GAI	-	: C A	A C f	et C	CC.		CAT	· C Δ .	A C C	·GT	cc/	A C A	4 T (CCTI	- "FO
G D K S V L A A K A K A A A S I V Y N N V A G S H A G T L G GCGGCGCCAGAGTGATAAGGGACCGTATTCGGCCATTGTCGGTATCAGCTTGGAGGATGGCCAGAAGCTGATCAAGCTTGCTGAGGCTGGA 63C A A Q S D K G P Y S A I V G I S L E D G Q K L I K L A E A G TCGGTATCTGTGGATCTGTGGGTGGATAGTAAGCAGGAGACCGTACGAGCGTATAACGTTGTCGCGCAGAAGCAAGGGCGGGATCCGAAC S V S V D L W V D S K Q E N R T T Y N V V A Q T K G G D P N AACGTCGTCGCGGCTGGGTGGCCACACGGACTCAGTCGAGGCGGGCCCTGGTATCAACGACGATGGCTCGGGCATTATTAGCAACTTGGTC 8100 N V V A L G G H T D S V E A G P G I N D D G S G I I S N L V ATTGCCCAAAGCGCTCACGCAGTACTCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAAGCAGGAGGAGTTCGGTCTGCTGGGCAGCACAAC POOL A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N TACCTACGTCTCCCATCTGAATGCCACCGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACATGATCGCCCTCACCTAACTACGCCCTC 990 Y Y V S H L N A T E L N K I R L Y L N F D M I A S P N Y A L ATTGCTATGACCGTGGTTGGATGGCGTTCAACCAGGAGCGGACCGGCCGG																																				
G D K S V L A A K A K A A A S I V Y N N V A G S H A G T L G GCGGCGCAGAGTGATAAGGGACCGTATTCGGCCATTGTCGGTATCAGCTTGGAGGATGGCCAGAAGCTGATCAAGCTTGCTGAGGCTGGA 63C A A Q S D K G P Y S A I V G I S L E D G Q K L I K L A E A G TCGGTATCTGTGGATCTGTGGGTTGGATGATGAAGCAGGAAACCGTACGAAGCTGTGCGGCAGAAGCGAAGGCGGGGATCCGAAC S V S V D L W V D S K Q E N R T T Y N V V A Q T K G G D P N AACGTCGTCGCGGCTGGGTGGCACACGGACTCAGTCGAGGCGGCCCTGGTATCAACGTTGTCGGCCAGAAGGAAG																					-	_	•				• .									
GEGGGCGAGAGTGATAAGGGACCGTATTCGGCCATTGTCGGTATCAGCTTGGAGGATGGCCAGAAGCTGATCAAGCTTGCTGAGGCTGGA 63C A A Q S D K G P Y S A I V G I S L E D G Q K L I K L A E A G TCGGTATCTGTGGATCTGTGGGTGGATAAGCAGGAGAACCGTACGACGTATAACGTTGTCGGCCAGACGAAGGGCGGGGGGGATCCGAAC 72C S V S V D L W V D S K Q E N R T T Y N V V A Q T K G G D P N AACGTCGTCGCGGCTGGGTGGCCACACGGACTCAGTCGAGGCGGGCCCTGGTATCAACGACGAGGATGGCTCGGCAATTATAGCAACTTGGTC 810 N V V A L G G H T D S V E A G P G I N D D G S G I I S N L V ATTGCCAAAGCGCTCACGCAGTACTCCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAGCAGAGGAGGAGTTCGGTCTGGGCAGCAAC 900 I A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N ACCTACGTCTCCCATCTGAATGCCACCAGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACAGTACTCGCCTCACCTAACTACGCCCTC 990 Y Y V S H L N A T E L N K I R L Y L N F D N I A S P N Y A L TGATCTATGACGGTGATGGCGTTCAACCAGAGACGGACCGGCCGG																																				
A A Q S O K G P Y S A I V G I S L E D G Q K L I K L A E A G CGGTATCTGTGGATCTGTGGGTGGATAGTAAGCAGGAGAACCGTACGACGTATAACGTTGTCGCCAGAGGAGGGCGGCGATCCGAAC S V S V D L W V D S K Q E N R T T Y N V V A Q T K G G D P N ACGTCGTCGCCGCTGGGTGGCCACACGGACTCAGTCGAGGGGGGCCCTGGTATCAACGACGATGGCTCGGGCATTATTAGCAACTTGGTC 810 N V V A L G G H T D S V E A G P G I N D D G S G I I S N L V TTGCCAAAGCGCTCACGCAGTACTCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAGCAGAGAGGAGTTCGGTCTGGCGAGCAAC 900 I A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N ACTACGTCTCCCATCTGAATGCCCACCGAGCTGAACAAGATCCGACTGACCTGACTTCGACATGATCGCCTCACCTAACTACGCCCTC 990 Y Y V S H L N A T E L N K I R L Y L N F D H I A S P N Y A L TGATCTATGACGGTGATGGATCGGCGTTCAACCAGGAGGGACCGGCCGG																							•													
TEGGTATCTGTGGATCTGTGGTGGATAGTAAGCAGGAAACCGTACGACGTATAACGTTGTCGCGCAGACGAAGGCGGCGATCCGAAC 720 S V S V D L W V D S K Q E N R T T Y N V V A Q T K G G D P N TAACGTCGTCGCGCTGGGTGGCCACACGGACTCAGTCGAGGCGGGCG																																				
S V S V D L W V D S K Q E N R T T Y N V V A Q T K G G D P N ACCGTCGTCGGCGGGGGGCCCACCGGACCCAGGCGGGCCCTGGTATCAACGACGATGGCTCGGGCATTATTAGCAACTTGGTC 810 N V V A L G G H T D S V E A G P G I N D D G S G I I S N L V TTGCCAAAGCGCTCACGCAGTACTCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAGCAAGAGAGTTCGGTCGG																																				
AACGTCGTCGCGCTGGGTGGCCACACGGACTCAGTCGAGGCGGGCCCTGGTATCAACGACGATGGCTCGGCCATTATTAGCAACTTGGTC 810 N V V A L G G H T D S V E A G P G I N D D G S G I I S N L V TTGGCCAAAGCGGTCACGCAGTACTCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAGCAGAGAGGATTCGGTCTGCTGGGCAGCAAC 900 I A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N ACTACGTCTCCCATCTGAATGCCACCGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACATGATCGCCTCAACTAACT																																	4			
N V V A L G G H T D S V E A G P G I N D D G S G I I S N L V TTGCCAAAGCGCTCACGCAGTACTCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAGCAGAGGAGTTCGGTCTGCTGGGCAGCAAC 900 I A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N ACTACGTCTCCCATCTGAATGCCACCGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACATGATCGCCCTCACCTAACTACGCCCTC 990 Y Y V S H L N A T E L N K I R L Y L N F D M I A S P N Y A L TGATCTATGACGGTGATGGATCGGCGTTCAACCAGAGCGGACCGGCCGG																																				
TTGCCAAAGCGCTCACGCAGTACTCCGTCAAGAATGCCGTGCGCTTCCTCTTCTGGACAGCAGAGGAGTTCGGTCTGCTGGGCAGCAAC 900 I A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N ACTACGTCTCCCATCTGAATGCCACCGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACATGATCGCCTCACCTAACTACGCCCTC 990 Y Y V S H L N A T E L N K I R L Y L N F D M I A S P N Y A L TGATCTATGACGGTGATGGATCGGCGTTCAACCAGAGCGGACCGGCCGG																																				810
I A K A L T Q Y S V K N A V R F L F W T A E E F G L L G S N ACTACGTCTCCCATCTGAATGCCACCGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACATGATCGCCTCACCTAACTACGCCCTC 990 Y Y V S H L N A T E L N K I R L Y L N F D M I A S P N Y A L FGATCTATGACGGTGATGGATCGGCGTTCAACCAGAGCGGACCGGCCGG																																				
ACTACGTCTCCCATCTGAATGCCACCGAGCTGAACAAGATCCGACTGTACCTGAACTTCGACATGATCGCCTCACCTAACTACGCCCTC Y Y V S H L N A T E L N K I R L Y L N F D M I A S P N Y A L TGATCTATGACGGTGATGGATCGGCGTTCAACCAGAGCGGACCGGCCGG																																				900
Y Y V S H L N A T E L N K I R L Y L N F D M I A S P N Y A L GATCTATGACGGTGATGGATCGGCGTTCAACCAGAGCGGACCGGCCGG	1	A	^	А	Ļ	1	(1	ř	5	٧	K	r	4	А	٧	ĸ	۲		L	F	W		ŀ	А	٤	٤	۲	į	à	L	L	G	S	N	
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CACGGGCGGAGGGCATCATGTCCGAAGAGAACGCAAGCCGCTGGGGAGGTCAAGCCGGCGTGGCCTACGACGCCAACTACCACGCC 1260 TO GOVANY HA GGGAGACAACATGACCAACCATGAAGCCTTCCTGATCAACTCCAAAGCCACCGCCTTCGCCGTCGCCACCTACGCCAACGAC 1350 GONMIN HEAFLINSKATAFAVATYAND CTCCTCGATCCCCAAACGGAATACCACATCCTTGCACCGACGACCACCGCCACCACGACCATCGGCAAGAGAGAG	4	1	Y	0	G	٥	G	i :	S	Α	F	N	a	1 :	S	G	٩	A	(G	S	Α	C	1	I	Ε	K	L	۶	•	Ε	0	Y	Y	0	
TCACGGGCGCGAGGGCATCATGTCCGAAGAGAACGCAAGCCGCTGGGGAGGTCAAGCCGGCGTGGCCTACGACGCCAACTACCACGCC 1260 F T G A E G I M S E E N A S R W G G O A G V A Y D A N Y H A GGGAGACAACATGACCAACCTCAACCATGAAGCCTTCCTGATCAACTCCAAAGCCACCGCCTTCGCCGTCGCCACCTACGCCAACGAC 1350 N G D N M T N L N H E A F L I N S K A T A F A V A T Y A N D GCTCCTCGATCCCCAAACGGAATACCACATCCTCCTTGCACCGACGACCACCGCCACCATGCGACCATTCGGCAAGAGAGACTCCGAAGACA 1440 S S I P K R N T T S S L H R R A R T M R P F G K R A P K T																																				1170
T G A E G I M S E E N A S R W G G O A G V A Y D A N Y H A GGGAGACAACATGACCATGAAGCCTTCCTGATCAACTCCAAAGCCACCGCCTTCGCCGTCGCCACCTACGCCAACGAC 1350 G D N M T N L N H E A F L I N S K A T A F A V A T Y A N D CTCCTCGATCCCCAAACGGAATACCACATCCTCCTTGCACCGACGAGCCCGCACCATGCGACCATTCGGCAAGAGAGACTCCGAAGACA 1440 S S I P K R N T T S S L H R R A R T M R P F G K R A P K T	i	I	0	L	P	Н	I	F		T	Q	F	0	(3	R	S	0	. Y	1	Ε	A	F	•	I	L	N	G	I	8	3	S	G	G	L	
GGGAGACAACATGACCAACCATGAAGCCTTCCTGATCAACTCCAAAGCCACCGCCTTCGCCGTCGCCACCTACGCCAACGAC 1350 G D N M T N L N H E A F L I N S K A T A F A V A T Y A N D CTCCTCGATCCCCAAACGGAATACCACATCCTCCTTGCACCGACGAGCCCGCACCATGCGACCATTCGGCAAGAGAGAG	CA	CGG	GCG	CG	GAG	GGC	ΑТ	CAT	GT	cce	SAAG	AG.	AAI	CGC	:AA	GCC	:GC	TGG	GG	AG	GT	CAA	.GC	CG	GCG	TGG	cc	TAC	GA	CGO	CA	AC1	FAC	CAC	GCC	1260
G D N M T N L N H E A F L I N S K A T A F A V A T Y A N D CTCCTCGATCCCCAAACGGAATACCACATCCTCCTTGCACCGACGAGCCCGCACCATGCGACCATTCGGCAAGAGAGACTCCGAAGACA 1440 S S I P K R N T T S S L H R R A R T M R P F G K R A P K T	•	T	G	Α	ε	G	ī	٢	t	S	ε	ε	N	Å	4	S	R	W	G	3	G	a	A	(3	٧	Α	Y	0	1	4	N	Y	Н	Α	
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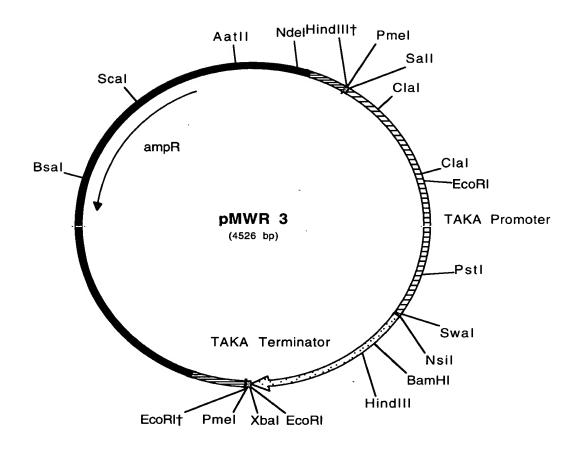


Fig. 2

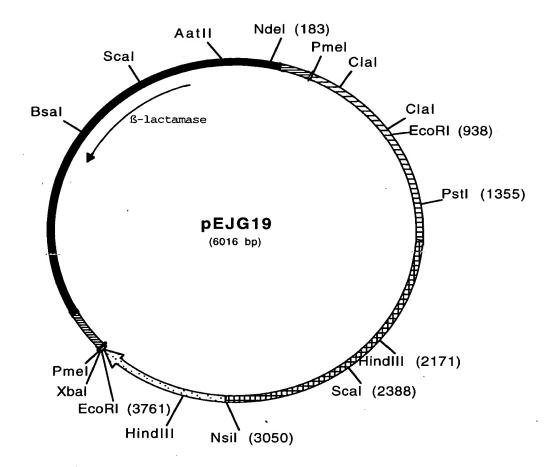


Fig. 3

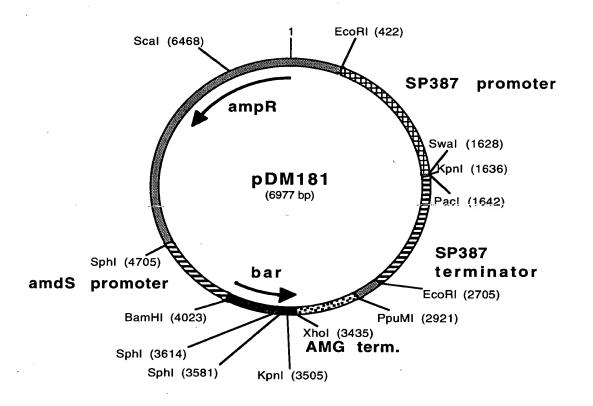


Fig. 4

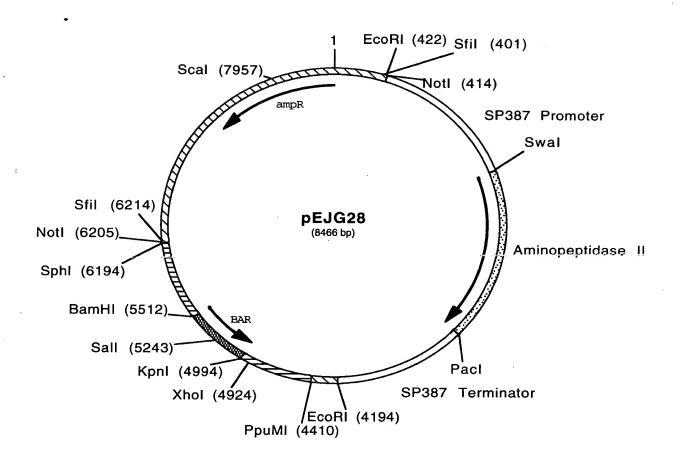


Fig. 5